OIPE

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/052,283

DATE: 02/12/2002
TIME: 12:01:23

Input Set : A:\P2751R1C1.txt
Output Set: N:\CRF3\02122002\J052283.raw

ENTERED

```
 3  <110> APPLICANT: Baker,Kevin P.
 4         Goddard,Audrey
 5         Wood,William I.
 7  <120> TITLE OF INVENTION: NOVEL POLYNUCLEOTIDES AND METHOD OF USE THEREOF
 9  <130> FILE REFERENCE: P2751R1C1
11  <140> CURRENT APPLICATION NUMBER: US/10/052,283
11  <141> CURRENT FILING DATE: 2002-01-15
11  <150> PRIOR APPLICATION NUMBER: PCT/US00/20006
12  <151> PRIOR FILING DATE: 2000-07-21
14  <150> PRIOR APPLICATION NUMBER: US 60/145,701
15  <151> PRIOR FILING DATE  1999-07-26
17  <160> NUMBER OF SEQ ID NOS: 564
19  <210> SEQ ID NO: 1
20  <211> LENGTH: 478
21  <212> TYPE: DNA
22  <213> ORGANISM: Homo Sapien
24  <220> FEATURE:
25  <221> NAME/KEY: unsure
26  <222> LOCATION: 212, 285, 424
27  <223> OTHER INFORMATION  unknown base
29  <400> SEQUENCE: 1
30    agtttgttaa aaataataat gccaataata tatgttattt tacgtatgtt 50
32    tatacagatg cacgcttatt tatacttatg tgtaagtgaa ataaatggca 100
34    aaaatgatac aaggcatagg aagaagaaat taggattata tgctatgtaa 150
36    gaagcagtat agtgtttttt gaaaatagac ttgaattagt tggaaatcca 200
38    tattgaaaac tntcgggcaa acattttaa aaaataaaaa aatgatatgc 250
40    taagaaagaa gagaaaacgg aattacacaa aatgntcaat taaaaccaca 300
42    aaaggaagca aaagtgtgga aaacaaaaag gggaacaaag aataaggcaa 350
44    caaacagaaa acagtaacaa atatggtaag cattaatcca actatattaa 400
46    taatcacttt aaatatcaat ggtntaaata tgtcaattat aagacagaga 450
48    ttaccagagt ggacacatta tataagct 478
50  <210> SEQ ID NO: 2
51  <211> LENGTH: 682
52  <212> TYPE: DNA
53  <213> ORGANISM: Homo Sapien
55  <220> FEATURE:
56  <221> NAME/KEY: unsure
57  <222> LOCATION: 45, 51, 62
58  <223> OTHER INFORMATION: unknown base
60  <400> SEQUENCE: 2
61    acgttcgcgg ttcgaggtta gggccccgga agcccgcagt aattncacgt 50
63    nttcccgccc gnttccggcc ccccagtgcc gcccttcgcg ggccggaggc 100
65    gcgagtctgg gctttggcgc cttcgcagcc gcaggcgaca tcctctttcc 150
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 67  tagctaaagc cccaaacgcc caggtggctt cctgggagag cacggctgag 200
 69  cctccgcctt cagatcagaa caggcagagc ctccaagggc ggcttgggcc 250
 71  cagtgcctgc ttatcctgcc cgtctctcca cacttcttct ttccctgtcc 300
 73  gttggagtcc attccttcct ggaaaagcca aagccgcgct cccctaagag 350
 75  tcatgtgtta ctggattaat tgaaattctt gataggtaac agagttttat 400
 77  catcagctta tgattgccta tgactagctc aaagttaagt tttaataaac 450
 79  tagtaagtac aataaacctc attcctaaat acaaggaaaa gaactatata 500
 81  atgaatactt gtctctatgc ccctctcgca tagataacaa taatttaggt 550
 83  ttacctttaa atgaactgca tttaaatgaa attaatttaa atgattgttt 600
 85  cacggcacag tttcatcaat ggtctacggt atcccttatt tatgtataca 650
 87  tcagtttgta tacatctgta tctatgtatt tg 682
 89  <210> SEQ ID NO: 3
 90  <211> LENGTH: 835
 91  <212> TYPE: DNA
 92  <213> ORGANISM: Homo Sapien
 94  <220> FEATURE:
 95  <221> NAME/KEY: unsure
 96  <222> LOCATION: 1, 6, 12, 27, 29-31, 49, 53, 56, 58, 61-62, 66, 68, 71, 74,
 97        78-81, 85-86, 88, 91, 95, 98, 102, 105, 109, 111, 115, 126-127,
 98        131, 139, 142, 146-147, 157, 163, 166, 169, 175, 177, 184,
 99        196-197, 199, 210, 215, 217, 219, 222, 228, 234, 236, 238,
100        240-241, 244-245, 257, 260, 263, 271, 279, 287, 291-331, 335,
101        337, 345, 384, 392, 418, 438, 459, 465, 478, 499, 501, 507, 533,
102        562, 579, 587, 598, 616, 630, 656, 661, 667, 671, 684, 702, 746,
103        811, 824-835
104  <223> OTHER INFORMATION: unknown base
106  <400> SEQUENCE: 3
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W--> 107  nagaanggaa gngaaggagg aagaggngnn naagagggag gggaaaagng 50
W--> 109  ganggngnag nngagnangg nggnggannn naggnngnag naggnccnag 100
W--> 111  anggnaagng nttgnaagaa agggannGcc nggtaaaana gnaccnnccc 150
W--> 113  aagaagngat tanggnggnt tcctngntga aggntgtgga tcccanntnt 200
W--> 115  ttcccgggan ttatngntng gnaacaanat ttcnangngn nacnnaggca 250
W--> 117  aacaatnaan ttnccaaggt nttggtagna tttcccncgg nnnnnnnnnn 300
W--> 119  nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn ngatntnggg gtttntcccc 350
W--> 121  tttcccctc ccctttcccc accccggggg ttcnggttgg tnaagaaaaa 400
W--> 123  aaaaaaaaaa aagaattntg gcgcggcctc ggcggagntg gtgatcggct 450
W--> 125  ggtgcatant cggcntctta ctactggnta ttttggcatt ctgctggana 500
W--> 127  natgttngta aataccaaag tcggcgggaa agngaagttg tttccaccat 550
W--> 129  aacagcaatt tnttttttag caattgcant tatcacntca gcacttgnac 600
W--> 131  cagaggaaat attttnggtt tcttacatgn aaaatcaaaa tggtacattt 650
W--> 133  aaggantggg ntaatgntaa ngtcagcaga cagnttgagg acactgtatt 700
W--> 135  anacggttac tatactttat attctgttat attgttctgt gtgttnttct 750
     137  ggatcccttt tgtctactta tattatgaag aaaaggatga tgatgatact 800
W--> 139  agtagatgta ntcaaattaa aacnnnnnnn nnnnn 835
     141  <210> SEQ ID NO: 4
     142  <211> LENGTH: 491
     143  <212> TYPE: DNA
     144  <213> ORGANISM: Homo Sapien
     146  <220> FEATURE:
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Input Set : A:\P2751R1C1.txt
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147  <221> NAME/KEY: unsure
148  <222> LOCATION: 181, 213, 251, 335, 345-346, 350, 357, 374, 381, 387, 398, 401,
149        403, 427, 442, 447, 451, 463, 481, 484, 486, 489
150  <223> OTHER INFORMATION: unknown base
152  <400> SEQUENCE: 4
153       agtttgttaa aaataataat gccaataata tatgttattt aacgtatgtt 50
155       tataacagat gcacgcttat ttatacttat gtgtaagtga aataaatggc 100
157       aaaaatgata caaggcatag gaagaagaaa ttaggattat atgctatgta 150
W--> 159  agaagcagta tagtgttttt tgaaaataga nttgaattag ttggaaatcc 200
W--> 161  atattgaaaa ctntcgggca aacattttta aaaaataaaa aaatgatatg 250
W--> 163  ntaagaaaga agagaaaacg gaattacaca aaatgctcaa ttaaaaccac 300
W--> 165  aaaaggaagc aaaagtgtgg aaaacaaaaa ggggnacaaa gaatnnggcn 350
W--> 167  acaaacngca aacagtaaca attntggtaa ncattantcc aattatantt 400
W--> 169  ncnattactc taaatatcaa tgttttnaat atgtctattg tnagacngag 450
W--> 171  nttaccagag agnacacatt atataaggtc ngangngtng g 491
173  <210> SEQ ID NO: 5
174  <211> LENGTH: 637
175  <212> TYPE: DNA
176  <213> ORGANISM: Homo Sapien
178  <220> FEATURE:
179  <221> NAME/KEY: unsure
180  <222> LOCATION: 4, 12-13, 31, 37, 40, 43-44, 46, 50, 55, 64, 68, 78, 97, 114,
181        124-125, 137, 148, 183, 185, 227, 233, 236, 238, 264, 266, 268,
182        276, 322, 325, 334, 340, 345, 417, 426, 428, 472, 474-475,
183        482-484, 500, 502-503, 509-510, 521, 540, 543, 558, 588-589, 594,
184        598, 606, 616, 631, 633
185  <223> OTHER INFORMATION: unknown base
187  <400> SEQUENCE: 5
W--> 188  ttcnttgtca anngtttttg gttcccccctt ntttccnggn ttnntntttn 50
W--> 190  ggaanaaaaa tttnaagnta taccaagnaa aaaattaaat tccaagnatt 100
W--> 192  ggattgaatt cccngggggat cttnnagaga tcccttngac tttgaccnaa 150
W--> 194  gggtccggct ttaggggaag aagttggtgt ttngntgggc cctggtactg 200
W--> 196  aagacgcgtt ccgggtagcc caaagangtt tcntantnac ccaaagcccc 250
W--> 198  gcacccgcct tttntntntt ttcttntggc aggatgaggc gtgcaggcct 300
W--> 200  gggtgaagga gtacttcctg gnaantatgg gaantatggn tatgntaata 350
     202  gtgggtatag tgcctgtgaa gaagaaaatg agaggctcac tgaaagtttg 400
W--> 204  agaagcaaag taactgntat aaaatntntt tcccattgaa ataggccatg 450
W--> 206  aagttaaaac ccagaataaa tnannagcgg annnggatta aaaagacgan 500
W--> 208  tnnacaacnn tgattttgta ngtataacta tgggcataan tgnagatttt 550
W--> 210  ttccagangg agctaaacaa agatgttgtg agatatgnng aggntatnat 600
W--> 212  taattntcaa gtttgntcac ataggcgagc ntnaaaac 637
214  <210> SEQ ID NO: 6
215  <211> LENGTH  969
216  <212> TYPE: DNA
217  <213> ORGANISM: Homo Sapien
219  <220> FEATURE:
220  <221> NAME/KEY: unsure
221  <222> LOCATION: 12-13, 25, 46, 49, 53, 64, 94, 118, 134, 140, 153, 155, 164, 172,
222        175, 177, 182, 184, 191, 210, 218, 221, 223-224, 240, 242, 250,
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Input Set   A:\P2751R1C1.txt
Output Set N:\CRF3\02122002\J052283.raw

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223        259, 266-267, 272, 276, 298, 305-306, 319, 339-340, 344, 346,
224        348, 353, 384, 397, 408, 432, 455, 457, 470, 474, 480, 502, 504,
225        519, 525, 559, 579, 597, 625, 629, 631, 634, 645, 649, 655, 657,
226        666, 670, 675, 681, 697, 768, 773, 776, 786, 788, 791, 794, 816,
227        819, 830, 859, 877, 879, 886, 895, 904, 910, 913, 918, 925,
228        927-946, 952, 956, 958, 962, 966, 968-969
229  <223> OTHER INFORMATION: unknown base
231  <400> SEQUENCE: 6
W--> 232  cccctttttc cnnggttttt ttttnggaaa aaaatttcag gggtanccng 50
W--> 234  ggnaaaattt aaantccagg gtttggggggg atttccccgg ggtncttttg 100
W--> 236  gagttccttt ggacctgnaa caaagggttg gaantaaaan aaaaattaaa 150
W--> 238  aancngggtt tttngggggaa anttnanaat gngnttgggg ncaagaaaaa 200
W--> 240  tgggtttttn gggagggnaa ngnnggttca tttccaaatn gnagggggggn 250
W--> 242  aaaaatttna ggcttnnggg gnaggnggaa aaaaatttcg tagcctcnag 300
W--> 244  gttgnnattt ttaaacctnc agaaggtggc cagccccgnn tcancngntg 350
W--> 246  atnaaggcag atgggaaaag ggggatatgg ggtnataagg gtacctntca 400
W--> 248  ccctttttnga aggaaaaaaa gtggtccaca gnatttttgt ttacccaagg 450
W--> 250  gtaananatg gaattttgtn gaanataggn gaatggtgag gcatttggaa 500
W--> 252  ananggggggg gggttttttnt tgaanggggg agtaggggta tggtatttta 550
W--> 254  tgggaaaana gttttttggc actaaaccnt tttgaattac ctaatanatt 600
W--> 256  tatgtggaaa cctgtccttt ttttncagnt naanaaaaat ttttncccnt 650
W--> 258  gaaantnatt tttagnaagn atatnaaaag nattttttttt ttcaagngtc 700
     260  agaaaccttt tagcatcatt gaagttaaaa tgactgtcca taaacttttc 750
W--> 262  agaaatagta ggcatttnag gcnacnagat ttgtanangg natnttcata 800
W--> 264  gaattatacc agtgganttna ccacctgaan cctcttggat cccgtaagca 850
W--> 266  ttctttgcna caaggaaggg aggtatncng ggtaantcct tgaanttttg 900
W--> 268  gacnggaacn atnacttnga atttnannnn nnnnnnnnnn nnnnnngccg 950
W--> 270  cnggggncntt tntcgngnn 969
     272  <210> SEQ ID NO: 7
     273  <211> LENGTH: 952
     274  <212> TYPE: DNA
     275  <213> ORGANISM: Homo Sapien
     277  <220> FEATURE:
     278  <221> NAME/KEY: unsure
     279  <222> LOCATION: 1, 3, 7, 23, 30, 38, 49, 53, 64, 68, 84, 90-92, 109, 130, 147,
     280        152, 182, 198, 211, 221-222, 228, 239, 243, 255, 271, 300, 303,
     281        308-309, 314, 322, 331, 337, 340, 345, 359, 377, 381, 385, 412,
     282        422, 452, 463, 469, 475, 541, 547, 586, 777, 780-781, 784-787,
     283        789, 798, 802-803, 825, 831, 838, 840, 853, 855, 863, 865, 872,
     284        885, 889-890, 907, 921-952
     285  <223> OTHER INFORMATION: unknown base
     287  <400> SEQUENCE: 7
W--> 288  ngntttngtt cccttttttc ccnggtttttn ttttggnaa aaaaatttna 50
W--> 290  ggnttaaccc aggnaaaanat taaattccaa gggntttggn nngaattccc 100
W--> 292  cggggggttnc ttttaggggt tcctttgacn ttgaaccaaa ggtttcnggc 150
W--> 294  cnggaggggg gggggaccgg tttttttcccc cngcgtttcc cccgggggntg 200
W--> 296  ggggttgggg ngcccatttg nngaagtnag tggggaggng gantgggaac 250
W--> 298  ccggnagttt tggagaaagg naggttcctt ccttaacccct gggggttccn 300
W--> 300  ggngcccnng gagnggcagt tngggggaata ntgtttnagn ggttnggggg 350
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```
W--> 302  gttttcctng ggtcccgcca aggggggnggt ncttnataaa agggtgcctt 400
W--> 304  tttccccaca gnttccaggt cngagaggag ccgcaccgtc gggttggaga 450
W--> 306  tngcgcgcaa ggnggcttnt ggttnggatt tgccccgcat cggccacagg 500
W--> 308  aaaagcctgg tccctaggca cggttgtggt tcgagctttt ngtttntcg 550
W--> 310  aacattgagg tattcgctca gcccaccacg ttgtcntcgg ggttattagg 600
     312  ccccagtcac aagccctatg atgttttcag acttcccagg tggagataag 650
     314  gaaaatttta ctatttctgc agaacttctg ttgatgtaca gcattgtatt 700
     316  tagcaacttc tgtgtagatc tgaaaataaa tacattacca attgttagtt 750
W--> 318  gcgtttttat taatataatt ttagagnagn ngannnngnt gttagacnta 800
W--> 320  cnnaggtaaa ttatgtggca ctttngcatt nttgttgntn catgttcccc 850
W--> 322  tgnantttgc ttngngattt cnatttattc caaantcann atagaatgta 900
W--> 324  atttccnaac ccacagtccg nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn 950
W--> 326  nn 952
     328  <210> SEQ ID NO: 8
     329  <211> LENGTH: 752
     330  <212> TYPE: DNA
     331  <213> ORGANISM  Homo Sapien
     333  <220> FEATURE:
     334  <221> NAME/KEY  unsure
     335  <222> LOCATION  4-6, 8, 11, 34, 37, 40, 87, 123, 125, 133-134, 138, 143, 147,
     336        166, 171, 174, 199, 242, 263, 302, 305, 333, 363, 380, 423, 425,
     337        463, 527, 534, 539, 547-548, 553, 556, 561, 583, 604, 607, 615,
     338        618, 621, 624, 628, 638, 641-643, 646, 648, 654-655, 658, 660,
     339        666, 674, 676, 679, 688-689, 704, 710-725, 727, 730, 735, 739,
     340        741, 745, 747, 749, 751-752
     341  <223> OTHER INFORMATION: unknown base
     343  <400> SEQUENCE  8
W--> 344  ggannngntt ncaaaatggg attttttaacc aaantanggn agagaaaagt 50
W--> 346  ttaagtgttt tgccaaaaaa attccaagga aaataangcg gagtttgatt 100
W--> 348  tttcagagtt caacaggaaa aangngaaca aanngccngg agntttnaaa 150
W--> 350  gttttgggaa agccantttt natntgttca aggaacagtt tttatttgng 200
W--> 352  atgccaatca gaattttgga cccagtataa tcaaggtcag antttcaacc 250
W--> 354  taagcctgga ccngacccat aataacggaa agtttaacaa tgactcacat 300
W--> 356  tntcntaaag tttccagcca gaataggaca cgntcatttg gtcattttcc 350
W--> 358  cggtccagag ttnttggatg tagagaaaan tagcttttcc caggaacaat 400
W--> 360  tttgtgattc cgcaggagaa ggntntgaaa gaatacatca agattttgaa 450
W--> 362  tttggtgatg aanttagcag cagctccact gaacagataa gggcaaccac 500
W--> 364  acctccaaat caaggaaggc cagattntcc tgtntatgnt aacctnnag 550
W--> 366  aantgnaaat ntcccagtat ggtcttcccc canttcttgg gagcctggta 600
W--> 368  attnagntta ttggngcntg nganactnat atagacanct nnnggngntg 650
W--> 370  ttannatnan cacagnggga catngnatng aagttggnna cctcttgctt 700
W--> 372  ggantcgggn nnnnnnnnnn nnnnncntcn ccgcngggnc ntttntngng 750
W--> 374  nn 752
     376  <210> SEQ ID NO: 9
     377  <211> LENGTH: 478
     378  <212> TYPE: DNA
     379  <213> ORGANISM  Homo Sapien
     381  <400> SEQUENCE  9
     382  agtttgttaa aaataataat gccaataata tatgttattt tacgtatgtt 50
```

VERIFICATION SUMMARY
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Input Set : A:\P2751R1C1.txt
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L:11 M:270 C  Current Application Number differs, Replaced Current Application No
L:11 M:271 C  Current Filing Date differs, Replaced Current Filing Date
L 38 M:341 W  (46) "n" or "Xaa" used, for SEQ ID#:1
L 40 M:341 W  (46) "n" or "Xaa" used, for SEQ ID#:1
L:46 M 341 W  (46) "n" or "Xaa" used, for SEQ ID#:1
L:61 M 341 W  (46) "n" or "Xaa" used, for SEQ ID#:2
L:63 M 341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:107 M 341 W  (46) "n" or "Xaa" used, for SEQ ID#:3
L:109 M:341 W  (46) "n" or "Xaa" used, for SEQ ID#:3
L:111 M 341 W  (46) "n" or "Xaa" used, for SEQ ID#:3
L.113 M 341 W  (46) "n" or "Xaa" used, for SEQ ID#:3
L 115 M 341 W  (46) "n" or "Xaa" used, for SEQ ID# 3
L:117 M:341 W  (46) "n" or "Xaa" used, for SEQ ID#:3
L:119 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L 121 M 341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:123 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:125 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:127 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L 129 M:341 W  (46) "n" or "Xaa" used, for SEQ ID#:3
L 131 M 341 W  (46) "n" or "Xaa" used, for SEQ ID# 3
L:133 M 341 W: (46) "n" or "Xaa" used, for SEQ ID# 3
L:135 M:341 W: (46) "n" or "Xaa" used, for SEQ ID# 3
L:139 M:341 W  (46) "n" or "Xaa" used, for SEQ ID# 3
L 159 M:341 W  (46) "n" or "Xaa" used, for SEQ ID#:4
L:161 M 341 W  (46) "n" or "Xaa" used, for SEQ ID#:4
L:163 M 341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:165 M 341 W: (46) "n" or "Xaa" used, for SEQ ID# 4
L:167 M 341 W: (46) "n" or "Xaa" used, for SEQ ID# 4
L:169 M 341 W: (46) "n" or "Xaa" used, for SEQ ID# 4
L:171 M 341 W: (46) "n" or "Xaa" used, for SEQ ID# 4
L:188 M:341 W  (46) "n" or "Xaa" used, for SEQ ID#:5
L 190 M 341 W  (46) "n" or "Xaa" used, for SEQ ID#:5
L 192 M:341 W  (46) "n" or "Xaa" used, for SEQ ID#:5
L:194 M:341 W  (46) "n" or "Xaa" used, for SEQ ID#:5
L:196 M:341 W  (46) "n" or "Xaa" used, for SEQ ID#:5
L:198 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:200 M:341 W  (46) "n" or "Xaa" used, for SEQ ID#:5
L:204 M:341 W  (46) "n" or "Xaa" used, for SEQ ID#:5
L:206 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:208 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:210 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:212 M 341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:232 M 341 W  (46) "n" or "Xaa" used, for SEQ ID#:6
L:234 M 341 W. (46) "n" or "Xaa" used, for SEQ ID#:6
L:236 M 341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:238 M 341 W  (46) "n" or "Xaa" used, for SEQ ID#:6
L 240 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:242 M 341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
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VERIFICATION SUMMARY
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```
L 244 M.341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L 246 M 341 W  (46) "n" or "Xaa" used, for SEQ ID#:6
L 248 M 341 W  (46) "n" or "Xaa" used, for SEQ ID#:6
L 250 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L.17046 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:564
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